

SEQUENCE LISTING

<110> Pharmacia & Upjohn

<120> CRYSTALLIZATION AND STRUCTURE DETERMINATION OF
STAPHYLOCOCCUS AUREUS
UDP-N-ACETYLENOLPYRUVYLGLUCOSAMINE REDUCTASE (S. aureus
MurB)

<130> 6241.NCP

<140> Unassigned

<141> 2000-08-04

<150> 60/147,164

<151> 1999-08-04

<160> 11

<170> PatentIn Ver. 2.1

<210> 1

<211> 326

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Recombinant
S. aureus MurB protein including polyhistidine
region

<400> 1

Met Arg Gly Ser His His His His His Thr Asp Pro Ile Asn Lys
1 5 10 15

Asp Ile Tyr Gln Ala Leu Gln Gln Leu Ile Pro Asn Glu Lys Ile Lys
20 25 30

Val Asp Glu Pro Leu Lys Arg Tyr Thr Tyr Thr Lys Thr Gly Gly Asn
35 40 45

Ala Asp Phe Tyr Ile Thr Pro Thr Lys Asn Glu Glu Val Gln Ala Val
50 55 60

Val Lys Tyr Ala Tyr Gln Asn Glu Ile Pro Val Thr Tyr Leu Gly Asn
65 70 75 80

Gly Ser Asn Ile Ile Ile Arg Glu Gly Gly Ile Arg Gly Ile Val Ile

100 110 120 130 140 150 160 170 180 190 200 210 220 230 240 250 260 270 280 290 300 310 320 325

85										90					95							
Ser	Leu	Leu	Ser	Leu	Asp	His	Ile	Glu	Val	Ser	Asp	Asp	Ala	Ile	Ile							
			100					105					110									
Ala	Gly	Ser	Gly	Ala	Ala	Ile	Ile	Asp	Val	Ser	Arg	Val	Ala	Arg	Asp							
		115					120					125										
Tyr	Ala	Leu	Thr	Gly	Leu	Glu	Phe	Ala	Cys	Gly	Ile	Pro	Gly	Ser	Ile							
	130					135					140											
Gly	Gly	Ala	Val	Tyr	Met	Asn	Ala	Gly	Ala	Tyr	Gly	Gly	Glu	Val	Lys							
145					150					155					160							
Asp	Cys	Ile	Asp	Tyr	Ala	Leu	Cys	Val	Asn	Glu	Gln	Gly	Ser	Leu	Ile							
			165						170					175								
Lys	Leu	Thr	Thr	Lys	Glu	Leu	Glu	Leu	Asp	Tyr	Arg	Asn	Ser	Ile	Ile							
			180					185					190									
Gln	Lys	Glu	His	Leu	Val	Val	Leu	Glu	Ala	Ala	Phe	Thr	Leu	Ala	Pro							
		195					200					205										
Gly	Lys	Met	Thr	Glu	Ile	Gln	Ala	Lys	Met	Asp	Asp	Leu	Thr	Glu	Arg							
	210					215					220											
Arg	Glu	Ser	Lys	Gln	Pro	Leu	Glu	Tyr	Pro	Ser	Cys	Gly	Ser	Val	Phe							
225					230					235					240							
Gln	Arg	Pro	Pro	Gly	His	Phe	Ala	Gly	Lys	Leu	Ile	Gln	Asp	Ser	Asn							
			245						250					255								
Leu	Gln	Gly	His	Arg	Ile	Gly	Gly	Val	Glu	Val	Ser	Thr	Lys	His	Ala							
		260						265					270									
Gly	Phe	Met	Val	Asn	Val	Asp	Asn	Gly	Thr	Ala	Thr	Asp	Tyr	Glu	Asn							
		275					280					285										
Leu	Ile	His	Tyr	Val	Gln	Lys	Thr	Val	Lys	Glu	Lys	Phe	Gly	Ile	Glu							
	290					295					300											
Leu	Asn	Arg	Glu	Val	Arg	Ile	Ile	Gly	Glu	His	Pro	Lys	Glu	Ser	Leu							
305					310					315					320							
Gln	Pro	Ser	Leu	Ile	Ser																	
			325																			

<210> 2
 <211> 342
 <212> PRT
 <213> Escherichia coli

<400> 2

Met	Asp	His	Ser	Leu	Lys	Pro	Trp	Asn	Thr	Phe	Gly	Ile	Asp	His	Asn
1				5					10					15	
Ala	Gln	His	Ile	Val	Cys	Ala	Glu	Asp	Glu	Gln	Gln	Leu	Leu	Asn	Ala
			20					25						30	
Trp	Gln	Tyr	Ala	Thr	Ala	Glu	Gly	Gln	Pro	Val	Leu	Ile	Leu	Gly	Glu
		35					40					45			
Gly	Ser	Asn	Val	Leu	Phe	Leu	Glu	Asp	Tyr	Arg	Gly	Thr	Val	Ile	Ile
	50					55					60				
Asn	Arg	Ile	Lys	Gly	Ile	Glu	Ile	His	Asp	Glu	Pro	Asp	Ala	Trp	Tyr
65					70					75					80
Leu	His	Val	Gly	Ala	Gly	Glu	Asn	Trp	His	Arg	Leu	Val	Lys	Tyr	Thr
			85						90					95	
Leu	Gln	Glu	Gly	Met	Pro	Gly	Leu	Glu	Asn	Leu	Ala	Leu	Ile	Pro	Gly
			100					105					110		
Cys	Val	Gly	Ser	Ser	Pro	Ile	Gln	Asn	Ile	Gly	Ala	Tyr	Gly	Val	Glu
		115					120					125			
Leu	Gln	Arg	Val	Cys	Ala	Tyr	Val	Asp	Ser	Val	Glu	Leu	Ala	Thr	Gly
		130				135					140				
Lys	Gln	Val	Arg	Leu	Thr	Ala	Lys	Glu	Cys	Arg	Phe	Gly	Tyr	Arg	Asp
145					150					155					160
Ser	Ile	Phe	Lys	His	Glu	Tyr	Gln	Asp	Arg	Phe	Ala	Ile	Val	Ala	Val
				165					170					175	
Gly	Leu	Arg	Leu	Pro	Lys	Glu	Trp	Gln	Pro	Val	Leu	Thr	Tyr	Gly	Asp
			180					185						190	
Leu	Thr	Arg	Leu	Asp	Pro	Thr	Thr	Val	Thr	Pro	Gln	Gln	Val	Phe	Asn
		195					200					205			
Ala	Val	Cys	His	Met	Arg	Thr	Thr	Lys	Leu	Pro	Asp	Pro	Lys	Val	Asn
	210					215					220				

Gly Asn Ala Gly Ser Phe Phe Lys Asn Pro Val Val Ser Ala Glu Thr
 225 230 235 240

Ala Lys Ala Leu Leu Ser Gln Phe Pro Thr Ala Pro Asn Tyr Pro Gln
 245 250 255

Ala Asp Gly Ser Val Lys Leu Ala Ala Gly Trp Leu Ile Asp Gln Cys
 260 265 270

Gln Leu Lys Gly Met Gln Ile Gly Gly Ala Ala Val His Arg Gln Gln
 275 280 285

Ala Leu Val Leu Ile Asn Glu Asp Asn Ala Lys Ser Glu Asp Val Val
 290 295 300

Gln Leu Ala His His Val Arg Gln Lys Val Gly Glu Lys Phe Asn Val
 305 310 315 320

Trp Leu Glu Pro Glu Val Arg Phe Ile Gly Ala Ser Gly Glu Val Ser
 325 330 335

Ala Val Glu Thr Ile Ser
 340

<210> 3

<211> 259

<212> PRT

<213> Helicobacter pylori

<400> 3

Met Leu Glu Thr Thr Ile Asp Phe Ser Arg Tyr Ser Ser Val Lys Ile
 1 5 10 15

Gly Thr Pro Leu Lys Val Ser Val Leu Glu Asn Asp Asp Glu Ile Ser
 20 25 30

Gln Glu His Gln Ile Ile Gly Leu Ala Asn Asn Leu Leu Ile Ala Pro
 35 40 45

Ser Ala Lys Asn Leu Ala Leu Leu Gly Lys Asn Tyr Asp Tyr Ile Cys
 50 55 60

Asp Lys Gly Glu Cys Val Glu Ile Gly Gly Ala Ala Asn Ala Ser Lys
 65 70 75 80

Ile Phe Asn Tyr Phe Arg Ala Asn Asp Leu Glu Gly Leu Glu Phe Leu
 85 90 95

Gly Gln Leu Pro Gly Thr Leu Gly Ala Leu Val Lys Met Asn Ala Gly
 100 105 110
 Met Lys Glu Phe Glu Ile Lys Asn Val Leu Glu Ser Ala Cys Ile Asn
 115 120 125
 Asn Gln Trp Leu Glu Lys Glu Ala Leu Gly Leu Gly Tyr Arg Ser Ser
 130 135 140
 Gly Phe Ser Gly Val Val Leu Arg Ala Arg Phe Lys Lys Thr His Gly
 145 150 155 160
 Phe Arg Glu Gly Val Leu Lys Ala Cys Gln Ser Met Arg Lys Ser His
 165 170 175
 Pro Lys Leu Pro Asn Phe Gly Ser Cys Phe Lys Asn Pro Pro Asn Asp
 180 185 190
 His Ala Gly Arg Leu Leu Glu Gly Val Gly Leu Arg Gly Tyr Cys Leu
 195 200 205
 Lys Arg Val Gly Phe Ala Lys Glu His Ala Asn Phe Leu Val Asn Leu
 210 215 220
 Gly Gly Ala Glu Phe Glu Glu Ala Leu Asp Leu Ile Glu Leu Ala Lys
 225 230 235 240
 Ala Arg Val Leu Gln Glu Tyr Gly Ile His Leu Glu Glu Glu Val Lys
 245 250 255
 Ile Leu Arg

<210> 4

<211> 297

<212> PRT

<213> Aquifex aeolicus

<400> 4

Met Leu Phe Leu Lys Asn Val Pro Leu Gln Asn Leu Thr Thr Ile Lys
 1 5 10 15

Ile Gly Gly Arg Val Ser Phe Tyr Ala Glu Pro Ser Asp Leu Lys Glu
 20 25 30

Ile Ser Leu Cys Ile Asp Phe Ser Lys Ser Arg Asp Ile Pro Leu Phe

35	40	45
Val Leu Gly Asn Gly Ser Asn Thr Ile Phe Gly Asp Val Arg Gly Leu		
50	55	60
Val Val Asn Leu Lys Asn Leu Lys Gly Phe Lys Val Lys Glu Ile Lys		
65	70	75 80
Gly Lys Phe Phe Val Glu Ala Phe Ser Gly Thr Pro Leu* Lys Asp Leu		
85	90	95
Ile Arg Phe Ser Val Lys Glu Asn Val Lys Ser Phe Tyr Lys Leu Leu		
100	105	110
Gly Phe Pro Ala Ser Val Gly Gly Ala Val Ser Met Asn Ala Gly Ala		
115	120	125
Phe Gly Val Glu Ile Ser Asp Phe Leu Lys Glu Val Tyr Phe Val Asp		
130	135	140
Trp Glu Gly Lys Leu Gln Lys Ala Lys Arg Asp Glu Leu Asn Phe Ser		
145	150	155 160
Tyr Arg Lys Ser Pro Phe Pro Lys Leu Gly Ile Val Phe Lys Val Val		
165	170	175
Phe Glu Phe Glu Arg Ser Lys Glu Asn Ile Leu Pro Lys Tyr Glu Lys		
180	185	190
Ile Arg Arg Ile Arg Lys Glu Lys Gln Pro Ile Asn Leu Pro Thr Ser		
195	200	205
Gly Ser Thr Phe Lys Asn Pro Glu Gly Asn Phe Ala Gly Lys Leu Leu		
210	215	220
Glu Lys Ala Gly Leu Lys Gly Phe Arg Leu Lys Asn Val Gly Phe Ser		
225	230	235 240
Glu Lys His Ala Asn Phe Leu Val Asn Tyr Gly Gly Gly Thr Phe Ser		
245	250	255
Glu Val Val Asp Leu Ile Asn Ile Ala Lys Glu Arg Val Tyr Glu Asn		
260	265	270
Phe Gly Ile Val Leu Glu Glu Glu Val Lys Leu Ile Glu Ser Ser Gly		
275	280	285
Ser Asp Gly Trp Lys Val Leu Gly Ala		

290

295

<210> 5

<211> 303

<212> PRT

<213> Bacillus subtilis

<400> 5

Met Glu Lys Val Ile Gln Glu Leu Lys Glu Arg Glu Val Gly Lys Val
 1 5 10 15

Leu Ala Asn Glu Pro Leu Ala Asn His Thr Thr Met Lys Ile Gly Gly
 20 25 30

Pro Ala Asp Val Leu Val Ile Pro Ser Ser Val Asp Ala Val Lys Asp
 35 40 45

Ile Met Asp Val Ile Lys Lys Tyr Asp Val Lys Trp Thr Val Ile Gly
 50 55 60

Arg Gly Ser Asn Leu Leu Val Leu Asp Glu Gly Ile Arg Gly Val Val
 65 70 75 80

Ile Lys Leu Gly Ala Gly Leu Asp His Leu Glu Leu Glu Gly Glu Gln
 85 90 95

Val Thr Val Gly Gly Gly Tyr Ser Val Val Arg Leu Ala Thr Ser Leu
 100 105 110

Ser Lys Lys Gly Leu Ser Gly Leu Glu Phe Ala Ala Gly Ile Pro Gly
 115 120 125

Ser Val Gly Gly Ala Val Tyr Met Asn Ala Gly Ala His Gly Ser Asp
 130 135 140

Met Ser Glu Ile Leu Val Lys Ala His Ile Leu Phe Glu Asp Gly Thr
 145 150 155 160

Ile Glu Trp Leu Thr Asn Glu Gln Met Asp Phe Ser Tyr Arg Thr Ser
 165 170 175

Val Leu Gln Lys Lys Arg Pro Gly Val Cys Leu Glu Ala Val Leu Gln
 180 185 190

Leu Glu Gln Lys Asp Lys Glu Ser Ile Val Gln Gln Met Gln Ser Asn
 195 200 205

Leu Gly Gly Ala Val Trp Met Asn Ala Arg Cys Phe Gly Asn Glu Ile
130 135 140

Ser Glu Ile Leu Lys Lys Ile Thr Phe Ile Asp Asp Lys Gly Lys Thr
145 150 155 160

Ile Cys Lys Glu Phe Lys Lys Glu Asp Phe Lys Tyr Lys Ile Ser Pro
165 170 175

Phe Gln Asn Lys Asn Phe Phe Ile Leu Lys Ile Glu Leu Asn Leu Lys
180 185 190

Lys Asp Asn Lys Lys Ile Ile Glu Glu Lys Met Asn Lys Asn Lys Gln
195 200 205

Ala Arg Ile Asn Arg Gly His Tyr Leu Phe Pro Ser Gly Gly Ser Thr
210 215 220

Phe Lys Asn Asn Lys Ala Phe Leu Lys Pro Ser Gly Gln Ile Ile Glu
225 230 235 240

Glu Cys Lys Leu Lys Gly Leu Ser Ile Gly Gly Ala Thr Val Ser Lys
245 250 255

Tyr His Gly Asn Phe Ile Ile Asn Ile Asn Asn Ala Thr Ser Lys Asp
260 265 270

Ile Lys Ser Leu Ile Glu Lys Val Lys Ala Glu Val Tyr Leu Lys Thr
275 280 285

Gly Leu Leu Leu Glu Glu Glu Val Leu Tyr Ile Gly Phe Lys
290 295 300

<210> 7

<211> 304

<212> PRT

<213> Chlamydia pneumoniae

<400> 7

Met Lys Glu Ala Ala Pro Met His Phe Pro Phe Pro Val Arg Arg Ser
1 5 10 15

Val Trp Leu Asn Arg Tyr Ser Thr Phe Arg Ile Gly Gly Pro Ala Asn
20 25 30

Tyr Phe Lys Ala Ile His Thr Ile Glu Glu Ala Arg Glu Val Ile Arg

35					40					45						
Phe	Leu	His	Ser	Ile	Asn	Tyr	Pro	Phe	Leu	Ile	Ile	Gly	Lys	Gly	Ser	
50					55					60						
Asn	Cys	Leu	Phe	Asp	Asp	Arg	Gly	Phe	Asp	Gly	Phe	Val	Leu	Tyr	Asn	
65					70					75					80	
Ala	Ile	Tyr	Gly	Lys	Gln	Phe	Leu	Glu	Asp	Ala	Arg	Ile	Lys	Ala	Tyr	
					85					90					95	
Ser	Gly	Leu	Ser	Phe	Ala	Ala	Leu	Gly	Lys	Ala	Thr	Ala	Tyr	Asn	Gly	
100					105					110						
Tyr	Ser	Gly	Leu	Glu	Phe	Ala	Ala	Gly	Ile	Pro	Gly	Ser	Val	Gly	Gly	
115					120					125						
Ala	Ile	Phe	Met	Asn	Ala	Gly	Thr	Asn	Glu	Ser	Asp	Ile	Ser	Ser	Val	
130					135					140						
Val	Arg	Asn	Val	Glu	Thr	Ile	Asn	Ser	Glu	Gly	Glu	Leu	Cys	Ser	Tyr	
145					150					155					160	
Ser	Val	Glu	Glu	Leu	Glu	Leu	Ser	Tyr	Arg	Ser	Ser	Arg	Phe	His	Arg	
165					170					175						
Gln	Gln	Glu	Phe	Ile	Leu	Ser	Ala	Thr	Phe	Gln	Leu	Ser	Lys	Lys	Gln	
180					185					190						
Val	Ser	Ala	Asp	His	Ser	Lys	Ser	Ile	Leu	Gln	His	Arg	Leu	Met	Thr	
195					200					205						
Gln	Pro	Tyr	Thr	Gln	Pro	Ser	Ala	Gly	Cys	Ile	Phe	Arg	Asn	Pro	Glu	
210					215					220						
Gly	Thr	Ser	Ala	Gly	Lys	Leu	Ile	Asp	Ala	Ala	Gly	Leu	Lys	Gly	Leu	
225					230					235					240	
Ala	Ile	Gly	Gly	Ala	Gln	Ile	Ser	Pro	Leu	His	Ala	Asn	Phe	Ile	Ile	
245					250					255						
Asn	Thr	Gly	Lys	Ala	Thr	Ser	Asp	Glu	Val	Lys	Gln	Leu	Ile	Ala	Ile	
260					265					270						
Ile	Gln	Ser	Thr	Leu	Lys	Thr	Gln	Gly	Ile	Asp	Leu	Glu	His	Glu	Ile	
275					280					285						
Arg	Ile	Ile	Pro	Tyr	Gln	Pro	Lys	Ile	His	Ser	Pro	Val	Ser	Glu	Lys	

290

295

300

<210> 8

<211> 310

<212> PRT

<213> Rickettsia prowazekii

<400> 8

Met Ile Gln Asn Pro Met Ile Lys Leu Cys Asn Glu Ser Asn Asn Met
 1 5 10 15

Ser Ile Leu Pro Ile Ile Lys Gly Glu Tyr Lys Lys Asp Tyr Asn Leu
 20 25 30

Lys His Leu Thr Trp Phe Lys Val Gly Gly Asn Ala Glu Ile Phe Phe
 35 40 45

Lys Pro Phe Asp Phe Ala Asp Leu Lys Ser Phe Leu Ile Gln Asn Lys
 50 55 60

Gln Lys Leu Pro Ile Thr Thr Phe Gly Ser Gly Ser Asn Ile Ile Ile
 65 70 75 80

Arg Asp Gly Gly Ile Glu Gly Val Val Ile Lys Leu Gly Gln Asn Phe
 85 90 95

Asn Lys Ile Glu Phe Leu Asp Asn His Leu Ile Val Gly Ser Ser Cys
 100 105 110

Leu Asn Tyr Asn Leu Ala Arg Phe Cys Gln Ala Asn Ala Ile Ser Gly
 115 120 125

Phe Glu Phe Leu Val Gly Ile Pro Gly Thr Ile Gly Gly Gly Val Ile
 130 135 140

Met Asn Ala Gly Ala Tyr Gly Ser Ala Phe Gln Asp Ile Ile Val Gln
 145 150 155 160

Val Glu Ala Leu Asp Phe Ser Gly Asn Phe Leu Thr Phe Thr Asn Lys
 165 170 175

Glu Ile Gly Phe Lys Tyr Arg Gly Asn Asn Leu Pro Lys Asp Leu Ile
 180 185 190

Asn Asn Gly Ile Tyr Gly Leu Glu Asn Leu Ala Leu Ile Pro Gly Cys
 100 105 110

Ala Gly Ser Ala Pro Ile Gln Asn Ile Gly Ala Tyr Gly Val Glu Phe
 115 120 125

Lys Asp Val Cys Asp Tyr Val Glu Val Leu Asn Leu Asn Thr Asn Glu
 130 135 140

Thr Phe Arg Leu Asp Thr Glu Gln Cys Glu Phe Gly Tyr Arg Glu Ser
 145 150 155 160

Ile Phe Lys His Arg Tyr Gln Gln Gly Tyr Val Ile Thr Ala Val Gly
 165 170 175

Leu Lys Leu Lys Lys Asp Trp Gln Pro Ile Leu Lys Tyr Gly Ser Leu
 180 185 190

Val Glu Phe Asp Pro Lys Thr Val Thr Ala Lys Gln Ile Phe Asp Glu
 195 200 205

Val Cys His Ile Arg Gln Ser Lys Leu Pro Asp Pro Asn Glu Val Gly
 210 215 220

Asn Ala Gly Ser Phe Phe Lys Asn Pro Val Val Ser Ser Glu His Phe
 225 230 235 240

Glu Glu Ile Lys Lys His His Glu Asn Leu Pro His Phe Pro Gln Ala
 245 250 255

Asp Gly Ser Val Lys Leu Ala Ala Gly Trp Leu Ile Asp Gln Cys Asn
 260 265 270

Leu Lys Gly Phe Gln Ile Gly Gly Ala Ala Val His Lys Lys Gln Ala
 275 280 285

Leu Val Leu Ile Asn Lys Asn Gly Ala Thr Gly Gln Asp Val Val Lys
 290 295 300

Leu Ala His His Val Arg Gln Thr Val Ala Glu Lys Phe Gly Val Tyr
 305 310 315 320

Leu Gln Pro Glu Val Arg Phe Ile Ser Ala Thr Gly Glu Val Asn Ser
 325 330 335

Glu Gln Ile Ile Thr
 340

Arg Leu Pro Gly Val Arg Leu Val Gly Gln Cys Ala Asp Ala Trp Val
85 90 95

Val Glu Ala Ala Ala Gly Glu Asn Trp His Gly Phe Val Thr Ala Cys
100 105 110

Val Asp Asn Gly Trp Asp Gly Leu Glu Asn Leu Ala Leu Ile Pro Gly
115 120 125

Thr Val Gly Ala Ala Pro Val Gln Asn Ile Gly Ala Tyr Gly Val Glu
130 135 140

Leu Ala Asp Arg Phe His Ser Leu Thr Ala Trp Asp Val Lys Gly Gly
145 150 155 160

Arg Trp Val Glu Met Gly Ala Ala Glu Cys Arg Phe Ala Tyr Arg Asp
165 170 175

Ser Phe Phe Lys His Gln Glu Pro Gly Ala Trp Val Ile Gly Ser Val
180 185 190

Arg Phe Ala Leu Pro Arg Pro Trp Gln Pro Val Leu Asp Tyr Pro Asp
195 200 205

Leu Gln Arg His Ala Ala Leu Asp Gly Ala Ala Pro Thr Ala Arg Ala
210 215 220

Val Tyr Asp Ala Val Cys Ala Ile Arg Arg Ala Lys Leu Pro Asp Pro
225 230 235 240

Ala Val Val Gly Asn Ala Gly Ser Phe Phe Lys Asn Pro Leu Val Asp
245 250 255

Ala Gly Thr Arg Gln Ala Leu Leu Gly Arg Phe Pro Gly Leu Val Ser
260 265 270

Tyr Pro Gln Pro Asp Gly Arg Tyr Lys Leu Ala Ala Gly Trp Leu Ile
275 280 285

Asp Gln Cys Gly Trp Lys Gly Arg Gln Leu Gly Ala Ala Gly Val His
290 295 300

Asp Arg Gln Ala Leu Val Leu Val Asn Arg Gly Gly Ala Gln Ala Arg
305 310 315 320

Asp Ile Met Ala Leu Ala Ala Ala Ile Gln Gly Asp Val Glu Arg Arg
325 330 335

Tyr Gly Val Arg Leu Glu Pro Glu Pro Val Val Val Pro Ala Arg
 340 345 350

Protein Data Bank